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1278 GAGAAAGTGGGGGGATGGCTAAGAAAGCTGGGAATAGGGAACAGAAGAGGGGTAGTGGGT 1219
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          241 GAGAAAGTGGGGGATGGCTAAGAAAGCTGGGAGATAGGGAACAGAAGAAGAGGGTAGTGGGTG 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 301 GGCTAGGGGGCTGCCTTATTTAAAGTGGTTGTTTATGATTCTTATACTAATTTATACAA 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     361 AGAIATIAAGGCCCIGTICAITAAGAAATIGITCCCTICCCCTGTGTTCAATGTITGTAA 420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 467; DB 4; Length 50000;
100.0%; Pred. No. 3.1e-223;
.ive 0; Mismatches 0; Indels 0;
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                                                                                                                                       GENERAL INTORNATION:
APPLICANT: Ryan, James W.
APPLICANT: Ryan, James W.
APPLICANT: Sprinkle, Terry Joe Curtis
APPLICANT: Venema, Richard C.
TITLE OF INVENTION: Human Aminopeptidase P Gene
FILE REFERENCE: MCG103
CURRENT APPLICATION NUMBER: US/09/146,053A
CURRENT FILING DATE: 1998-09-02
EARLIER APPLICATION NUMBER: 60/057,854
EARLIER FILING DATE: 1997-09-02
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin Ver. 2.0
ALIGNMENTS
                                                                             3-09-146-053-4/c
Sequence 4, Application US/09146053A
Patent No. 6399349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 467; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
CORGANISM: Homo sapiens
US-09-146-053-4
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RESULT 2
US-08-863-639A-31/C
US-08-863-639A-31/C
Requence 31, Application US/08863639A
Ret. INFORMATION:
CAL INFORMATION:
TO Coassin, Peter J.
Rampal, Jang B.
The Cassin, Jang B.
The Cas

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Gaps
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GRNERAL INFORMATION:
GERSHON, Elliot S.
Badner, Judith A.
Goldin, Lynn R.
Berrettini, Wade H.
Yoshikawa, Takeo
Sanders, Alan R.
Esterling, Liss R.
Esterling, Liss C.
TITLE OF INVENTION: Chromosomal Markers and Diagnostic
TITLE OF INVENTION: Chromosomal Markers and Diagnostic
Tests for Manic-Depressive Illness
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5.4%; Score 25; DB 2; Length 36;
100.0%; Pred. No. 0.0046;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                  COMPUTER KALLANDLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage COMPUTER: IBM compatible operation SYSTEM: Windows 95 SOFTWARE: Corel WordPerfect 8 version CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/863,639A FILING DATE: May 28, 1997

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Joseph E. Muech Reference/DOCKET NUMBER: 20,532

REFERENCE/DOCKET NUMBER: 20,532

REFERENCE/DOCKET NUMBER: 11859-1

TELEPHONE: (626) 796-4000

TELEPHONE: (626) 796-4000

TELEPHONE: (626) 796-5321

INFORMATION FOR SEQ ID NO: 31: FURNAMENT CHERENCE/STERISTICS: TOWNEY: TEMPORMATICN: TELEPHONE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE PORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/091,952A
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ADDRESSEE: Townsend and Townsend and Crew LLi
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
FITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS
                         NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abeldon & Mak
STREET: 225 South Lake Avenue, 9th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          53 CGCIGCIGCIGCIGCIGCIGCI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: Other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 100.
Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 36 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     single
                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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                                                                                                                        Pasadena
                                                                                                                                                                      USA
                                                                                                                                                                      COUNTRY: US
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US-09-091-952A-15
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